

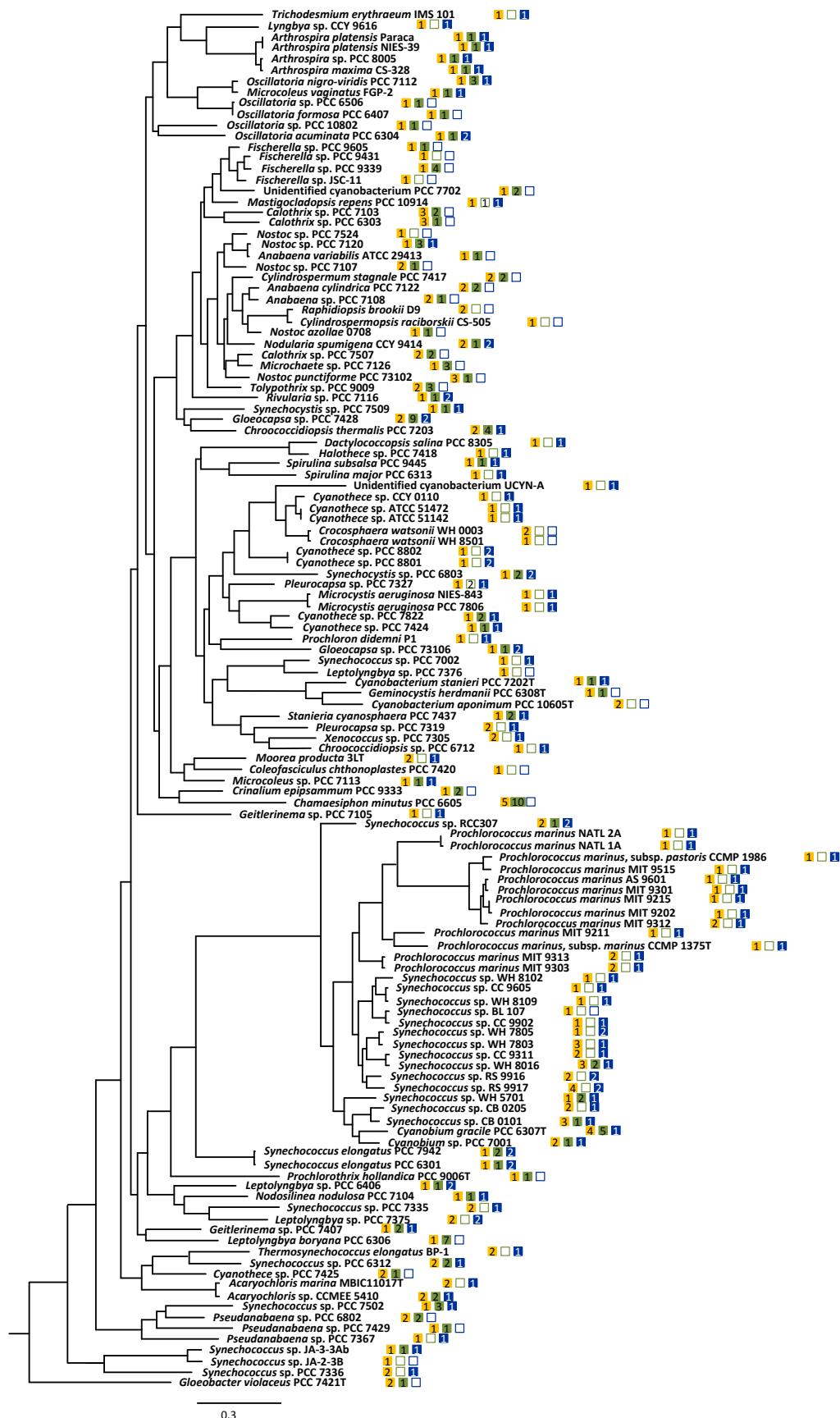
Supplementary Material

Extracellular proteins: Novel key components of metal homeostasis in cyanobacteria?

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Supplementary Figure 1. Distribution of TolC (yellow box), CopM (green box) and FutA2 (blue box) within the cyanobacterial phylum. The number of proteins containing the typical domains in the theoretical proteome of each strain is indicated within the box. Empty boxes correspond to the absence of CopM and/or FutA2. The cyanobacterial phylogenetic tree was generated by the maximum likelihood analysis of 31 conserved proteins concatenated by Shih et al., 2013. The proteins TolC, CopM and FutA2 of *Synechocystis* sp. PCC6803 were screened for their Pfam domains using available tools in the Integrated Microbial Genomes (IMG) database (v4.530, June 2015 <https://img.jgi.doe.gov/cgi-bin/m/main.cgi>). The presence or absence of genes encoding proteins with the selected Pfam domains was investigated in 126 cyanobacterial genomes, also available at the IMG. The list of encoding genes was refined in tblastn searches using the *Synechocystis* sp. PCC6803 sequences as query.



References

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